This listing of claims will replace all prior versions and listings of claims in the application:

## **Listing of Claims:**

- 1. (original) An isolated polypeptide comprising a BOG polypeptide fragment, said BOG fragment comprising a pRb binding motif and a casein kinase II phosphorylation motif.
- 2. (original) The BOG polypeptide fragment of claim 1, wherein said BOG polypeptide fragment is a full length BOG polypeptide.
- 3. (original) The BOG polypeptide fragment of claim 1, comprising an amino acid sequence as shown in Table 1, 5 or 7.
- 4. (original) The BOG polypeptide fragment of claim 1, wherein said casein kinase II phosphorylation motif is located downstream of the pRb binding motif.
- 5. (original) The BOG polypeptide fragment of claim 4, further comprising a second casein kinase II phosphorylation motif, said second casein kinase II phosphorylation motif being located upstream of the pRb binding motif.
  - 6. (original) The BOG polypeptide fragment of claim 1 joined to a detectable label.
- 7. (original) The BOG polypeptide fragment of claim 6, wherein the detectable label includes a radioactive isotope, an enzyme, a chromophore or a mixture thereof.
  - 8-22. (cancelled)
- 23. (original) A chimeric molecule comprising a BOG polypeptide fragment fused to a heterologous amino acid sequence.
  - 24-32. (cancelled)

- 33. (new) An isolated polypeptide comprising:
- i) at least 90% amino acid sequence identity with SEQ ID NO:2, SEQ ID NO:8 or SEQ ID NO:10;
- ii) a retinoblastoma gene produce (pRB) binding motif; and
- iii) at least one casein kinase II phosphorylation motif; wherein the polypeptide binds pRB and displaces E2F-1 bound to pRB.
- 34. (new) The polypeptide of claim 33 comprising an amino acid sequence of SEQ ID NO:2.
- 35. (new) The polypeptide of claim 33 comprising an amino acid sequence of SEQ ID NO:8.
- 36. (new) The polypeptide of claim 33 comprising an amino acid sequence of SEQ ID NO:10.
- 37. (new) The polypeptide of claim 33, wherein said casein kinase II phosphorylation motif is located downstream of the pRB binding motif.
- 38. (new) The polypeptide of claim 37 further comprising a second case II phosphorylation motif, said second case in kinase II phosphorylation motif being located upstream of the pRB binding motif.
  - 39. (new) The polypeptide of claim 33 joined to a detectable label.
- 40. (new) The polypeptide of claim 39, wherein the detectable label comprises a radioactive isotope, an enzyme, a chromophore or a mixture thereof.
- 41. (new) The polypeptide of claim 33 further comprising a heterologous amino acid sequence.

- 42. (new) The polypeptide of claim 41, wherein the heterologous amino acid is a tag polypeptide.
- 43. (new) The polypeptide of claim 41, wherein the heterologous amino acid sequence is that of an immunoglobulin constant region.
- 44. (new) The polypeptide of claim 41, wherein the heterologous amino acid sequence is maltose binding protein.